Ncol RFLP of the human LHRH gene on chromosome 8p

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Source and Description of Clone: A 0.2 kb EcoRI-SacI fragment from human LHRH cDNA cloned into SP64.

Polymorphism: NcoI identifies two alleles: Allele 1 — absence of 5.5 kb fragment Allele 2 — presence of 5.5 kb fragment Constant bands also at 9.5 and 3.2 kb.

Frequencies: 55/62 unrelated Caucasians were of the 1, 1 genotype. Observed heterozygosity = 0.11.

Assuming Hardy-Weinberg equilibrium:

Allele A1 (-5.5 kb): 0.94 Allele A2 (+5.5 kb): 0.06

Not Polymorphic For: PstI, TaqI, EcoRI, SstI, BamHI, BglIII, PvuII, MboI, EcoRV, HindIII, ApaI, BstNI, BcII, BgII, BstEII, BstNI, BstXI, HincII, HinfI, HphI, KpnI, StuI, XbaI and XmnI in 9 unrelated Caucasians.

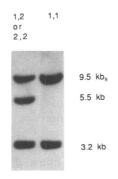
Chromosomal Localisation: The LHRH gene has been mapped to 8p21-p11.2 (2).

Mendelian Inheritance: Mendelian inheritance was demonstrated in four two-generation families (8 meioses scored).

Probe Availability: Contact A.J. Mason at Genentech, Inc.

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References: 1) J.P.Adelman et al. (1986) Proc. Natl. Acad. Sci. USA 83, 179-183. 2) T.L. Yang-Feng et al. (1986) Somatic Cell Molec. Genet. 12, 95-100.



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Source/Description: The probe pKSR2 is a 507 bp genomic fragment cloned into pUC19 vector. The pKSR2 was found during the sequence analyses of the cloned human endogenous HTLV-1 related genomic DNA (1).

Polymorphism:

Enzyme	Allele Size	Frequency*
<i>Eco</i> RI	A1 17.0 kb	0.55
	A2 3.2 kb	0.45
Observed het	erozygosity = 0.37	7
<i>Taq</i> I	B1 3.3 kb	0.25
	B2 1.9 kb	0.75

Observed heterozygosity = 0.26

*Studied in 19 unrelated Japanese individuals.

No Polymorphism Observed For: AluI, HaeIII, MspI, PvuII, RsaI (19 unrelated Japanese individuals).

Mendelian Inheritance: Co-dominant segregation was observed in 2 informative families.

Chromosomal Localization: Assigned to chromosome 8pter-8p22 using somatic cell hybrids (2).

Probe Availability: Contact Kiyotoshi Kaneko at the above address.

References: 1) Kaneko, K., Sato, S., Miyatake, T. and Tsuji, S. (1991) Neurology 41, 31-34. 2) Wagner, M.J., Ge, Y., Siciliano, M. and Wells, D.E. (1991) Genomics 10, 114-125.

A new RFLP locus D8S163 maps to human chromosome 8pter-8p22

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